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## SEQUENCE LISTING

<110> Japan Science and Technology Agency National Institute of Advanced Industrial Science and Technology
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<150> JP 2004-2192 <151> 2004-01-07
<150> JP 2004–93796 <151> 2004–03–26
<150> JP 2004–221592 <151> 2004–07–29
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Asp His Asp Tyr Leu Leu Gly Phe Ser Trp Pro Pro Arg Ser Tyr Thr
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Cys Ser Phe Cys Lys Arg Glu Phe Arg Ser Ala Gln Ala Leu Gly Gly 50 55 60

His Met Asn Val His Arg Arg Asp Arg Ala Arg Leu Arg Leu Gln Gln 65 70 75 80

Ser Pro Ser Ser Ser Ser Thr Pro Ser Pro Pro Tyr Pro Asn Pro Asn 85 90 95

Tyr Ser Tyr Ser Thr Met Ala Asn Ser Pro Pro Pro His His Ser Pro 100 105 110

Leu Thr Leu Phe Pro Thr Leu Ser Pro Pro Ser Ser Pro Arg Tyr Arg 115 120 125

Ala Gly Leu Ile Arg Ser Leu Ser Pro Lys Ser Lys His Thr Pro Glu 130 135 140

Asn Ala Cys Lys Thr Lys Lys Ser Ser Leu Leu Val Glu Ala Gly Glu 145 150 155 160

Ala Thr Arg Phe Thr Ser Lys Asp Ala Cys Lys IIe Leu Arg Asn Asp 165 170 175

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# Synthesized DNA Sequence

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## Synthesized DNA Sequence

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\223/	Synthesized DNA Sequence		
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caaac	ggagt cgtagatc		18
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#### Synthesized DNA Sequence

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His Glu Leu Thr Val Leu Cys Asp Ala Arg Val Ser Ile Ile Met Phe
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Ser Ser Ser Asn Lys Leu His Glu Tyr lle Ser Pro Asn Thr Thr
Lys Glu lle Val Asp Leu Tyr Gln Thr lle Ser Asp Val Asp Val Trp
                    70
                                        75
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                                     90
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                            120
Glu Asn Thr Phe Lys Leu Val Arg Glu Arg Lys Phe Lys Ser Leu Gly
                        135
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Asn Gln Ile Glu Thr Thr Lys Lys Lys Asn Lys Ser Gln Gln Asp Ile
145
                    150
Gln Lys Asn Leu 11e His Glu Leu Glu Leu Arg Ala Glu Asp Pro His
                165
                                    170
Tyr Gly Leu Val Asp Asn Gly Gly Asp Tyr Asp Ser Val Leu Gly Tyr
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Gln lle Glu Gly Ser Arg Ala Tyr Ala Leu Arg Phe His Gln Asn His
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His His Tyr Tyr Pro Asn His Gly Leu His Ala Pro Ser Ala Ser Asp
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gctagggttt cgattatcat gttctctagc tccaacaagc ttcatgagta tatcagccct 180
aacaccacaa cgaaggagat cgtagatctg taccaaacta tttctgatgt cgatgtttgg
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gccactcaat atgagcgaat gcaagaaacc aagaggaaac tgttggagac aaatagaaat
ctccggactc agatcaagca gaggctaggt gagtgtttgg acgagcttga cattcaggag 360
ctgcgtcgtc ttgaggatga aatggaaaac actttcaaac tcgttcgcga gcgcaagttc 420
aaatctcttg ggaatcagat cgagaccacc aagaaaaaga acaaaagtca acaagacata 480
caaaagaatc tcatacatga gctggaacta agagctgaag atcctcacta tggactagta
gacaatggag gagattacga ctcagttctt ggataccaaa tcgaagggtc acgtgcttac
                                                                   600
                                                                  660
gctcttcgtt tccaccagaa ccatcaccac tattacccca accatggcct tcatgcaccc
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<211> 365

<212> PRT

<213> Arabidopsis thaliana

<400> 136

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Tyr	Leu	Arg 35	Lys	Lys	Val	Asn	Ser 40	He	Glu	lle	Asp	Leu 45	Asp	Val	He
Arg	Asp 50	Val	Asp	Leu	Asn	Lys 55	Leu	Glu	Pro	Trp	Asp 60	lle	Gln	Glu	Met
Cys 65	Lys	lle	Gly	Thr	Thr 70	Pro	Gln	Asn	Asp	Trp 75	Tyr	Phe	Phe	Ser	His 80
Lys	Asp	Lys	Lys	Tyr 85	Pro	Thr	Gly	Thr	Arg 90	Thr	Asn	Arg	Ala	Thr 95	Ala
Ala	Gly	Phe	Trp 100	Lys	Ala	Thr	Gly	Arg 105	Asp	Lys	lle	lle	Tyr 110	Ser	Asn
Gly	Arg	Arg 115	lle	Gly	Met	Arg	Lys 120	Thr	Leu	Val	Phe	Tyr 125	Lys	Gly	Arg
Ala	Pro 130	His	Gly	GIn	Lys	Ser 135	Asp	Trp	He	Met	His 140	Glu	Tyr	Arg	Leu
Asp 145	Asp	Asn	lle	He	Ser 150	Pro	Glu	Asp	Val	Thr 155		His	Glu	Val	Val 160
Ser	He	He	Gly	Glu 165	Ala	Ser	Gln	Asp	Glu 170	Gly	Trp	Val	Val	Cys 175	Arg
He	Phe	Lys	Lys 180	Lys	Asn	Leu	His	Lys 185	Thr	Leu	Asn	Ser	Pro 190	Val	Gly
Gly	Ala	Ser 195	Leu	Ser	Gly	Gly	Gly 200	Asp	Thr	Pro	Lys	Thr 205	Thr	Ser	Ser
GIn	11e 210	Phe	Asn	Glu	Asp	Thr 215	Leu	Asp	Gin	Phe	Leu 220	Glu	Leu	Met	Gly
Arg 225	Ser	Cys	Lys	Glu	Glu 230	Leu	Asn	Leu	Asp	Pro 235	Phe	Met	Lys	Leu	Pro 240
Asn	Leu	Glu	Ser	Pro 245	Asn	Ser	GIn	Ala	11e 250	Asn	Asn	Cys	His	Va I 255	Ser
Ser	Pro	Asp	Thr	Asn	His	Asn	He	His	Val	Ser	Asn	Val	Val	Asp	Thr

260 265 270

Ser Phe Val Thr Ser Trp Ala Ala Leu Asp Arg Leu Val Ala Ser Gln 275 280 285

Leu Asn Gly Pro Thr Ser Tyr Ser IIe Thr Ala Val Asn Glu Ser His 290 295 300

Val Gly His Asp His Leu Ala Leu Pro Ser Val Arg Ser Pro Tyr Pro 305 310 315 320

Ser Leu Asn Arg Ser Ala Ser Tyr His Ala Gly Leu Thr Gln Glu Tyr 325 330 335

Thr Pro Glu Met Glu Leu Trp Asn Thr Thr Thr Ser Ser Leu Ser Ser 340 350

Ser Pro Gly Pro Phe Cys His Val Ser Asn Gly Ser Gly 355 360 365

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<212> DNA

<213> Arabidopsis thaliana

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His IIe His Gly Phe Cys Leu Gln Arg Cys Gly Lys Ser Cys Arg Leu 50 55 60

Arg Trp IIe Asn Tyr Leu Arg Pro Asp Leu Lys Arg Gly Ser Phe Ser 65 70 75 80

Pro Gln Glu Ala Ala Leu IIe IIe Glu Leu His Ser IIe Leu Gly Asn 85 90 95

Arg Trp Ala Gln IIe Ala Lys His Leu Pro Gly Arg Thr Asp Asn Glu 100 105 110

Val Lys Asn Phe Trp Asn Ser Ser IIe Lys Lys Lys Leu Met Ser His 115 120 125

His His His Gly His His His His Leu Ser Ser Met Ala Ser Leu 130 135 140

Leu Thr Asn Leu Pro Tyr His Asn Gly Phe Asn Pro Thr Thr Val Asp 145 150 155 160

Asp Glu Ser Ser Arg Phe Met Ser Asn IIe IIe Thr Asn Thr Asn Pro

Asn Phe IIe Thr Pro Ser His Leu Ser Leu Pro Ser Pro His Val Met 180 185 190

Thr Pro Leu Met Phe Pro Thr Ser Arg Glu Gly Asp Phe Lys Phe Leu 195 200 205

Thr Thr Asn Asn Pro Asn Gln Ser His His His Asp Asn Asn His Tyr 210 215 220

Asn Asn Leu Asp IIe Leu Ser Pro Thr Pro Thr IIe Asn Asn His His 225 230 235 240 Gln Pro Ser Leu Ser Ser Cys Pro His Asp Asn Asn Leu Gln Trp Pro 245 Ala Leu Pro Asp Phe Pro Ala Ser Thr Ile Ser Gly Phe Gin Glu Thr 265 Leu Gin Asp Tyr Asp Asp Ala Asn Lys Leu Asn Val Phe Val Thr Pro 285 275 280 Phe Asn Asp Asn Ala Lys Lys Leu Leu Cys Gly Glu Val Leu Glu Gly 295 Lys Val Leu Ser Ser Ser Pro Ile Ser Gln Asp His Gly Leu Phe 310 315 Leu Pro Thr Thr Tyr Asn Phe Gln Met Thr Ser Thr Ser Asp His Gln 325 330 335 His His His Arg Val Asp Ser Tyr Ile Asn His Met Ile Ile Pro Ser 345 350 340 Ser Ser Ser Ser Pro IIe Ser Cys Gly Gln Tyr Val IIe Thr 360 <210> 139 <211> 1104 <212> DNA

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Ala Tyr Glu Leu Ser Val Leu Cys Asp Ala Glu Val Ala Leu IIe Val 50 55 60

Phe Ser Ser Arg Gly Arg Leu Tyr Glu Tyr Ser Asn Asn Ser Val Lys 65 70 75 80

Gly Thr lle Glu Arg Tyr Lys Lys Ala lle Ser Asp Asn Ser Asn Thr 85 90 95

Gly Ser Val Ala Glu lle Asn Ala Gln Tyr Tyr Gln Gln Glu Ser Ala 100 105 110

Lys Leu Arg Gln Gln IIe IIe Ser IIe Gln Asn Ser Asn Arg Gln Leu 115 120 125

Met Gly Glu Thr lle Gly Ser Met Ser Pro Lys Glu Leu Arg Asn Leu 130 135 140

Glu Gly Arg Leu Glu Arg Ser lle Thr Arg lle Arg Ser Lys Lys Asn 145 150 155 160

Glu Leu Leu Phe Ser Glu l'Ie Asp Tyr Met Gln Lys Arg Glu Val Asp 165 170 175

Leu His Asn Asp Asn Gln IIe Leu Arg Ala Lys IIe Ala Glu Asn Glu 180 185 190

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Gin Leu Met Pro Pro Pro Gin Thr Gin Ser Gin Pro Phe Asp Ser Arg
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Asn Tyr Phe Gln Val Ala Ala Leu Gln Pro Asn Asn His His Tyr Ser
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gaaattaatg cacagtatta tcaacaagaa tcagccaaat tgcgtcaaca aataatcagc 360
atacaaaact ccaacaggca attgatgggt gagacgatag ggtcaatgtc tcccaaagag 420
ctcaggaact tggaaggcag attagagaga agtattaccc gaatccgatc caagaagaat 480
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